



Corrigendum: Flux balance analysis reveals acetate metabolism modulates cyclic electron flow and alternative glycolytic pathways in *Chlamydomonas reinhardtii*

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Reason for Corrigendum:

There was a mistake with the naming of the genome-scale metabolic model of *Chlamydomonas reinhardtii* originally published in *Molecular Systems Biology* in 2011 (Chang et al., 2011). In our article there are 10 instances where we refer to the model as *iCR1080*, however the correct name should be *iRC1080*, named after the initials of the researcher who reconstructed it. The authors apologize for the mistake. This error does not affect the scientific conclusions of the article in any way.

AUTHOR CONTRIBUTIONS

All authors listed have made substantial, direct and intellectual contribution to the work, and approved it for publication.

REFERENCES

Chang, R. L., Ghamsari, L., Manichaikul, A., Hom, E. F., Balaji, S., Fu, W., et al. (2011). Metabolic network reconstruction of *Chlamydomonas* offers insight into light-driven algal metabolism. *Mol. Syst. Biol.* 7, 52. doi: 10.1038/msb.2011.52

Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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